

# SEQUENCE LISTING

5 <110> McKeehan, Wallace L.  
 Luo, Yongde  
 <120> FGF-Affinity Chromatography  
 <130> 12740.0232.NPUS00 TAMK:232  
 10 <150> US 60/325,613  
 <151> 2001-09-28  
 <150> US 60/325,502  
 <151> 2001-08-28  
 15 <150> US 60/277,735  
 <151> 2001-03-21  
 20 <150> US 60/252,225  
 <151> 2000-11-21  
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 <170> PatentIn version 3.1  
 25 <210> 1  
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 Met Arg Lys  
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 165  
 Trp Ile Leu Thr Arg Ile Leu Pro Thr Pro Leu Tyr Arg Pro Cys Phe  
 5 10 15  
 50 cac ctc gtc tgt ctt gtg ggc acc ata tct tta gct tgc aat gac atg  
 213  
 His Leu Val Cys Leu Val Gly Thr Ile Ser Leu Ala Cys Asn Asp Met  
 20 25 30 35  
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Ser Pro Glu Gln Thr Ala Thr Ser Val Asn Cys Ser Ser Pro Glu Arg  
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10 aga ctg ttc tgt cgc acc cag tgg tac ctg agg att gac aaa cga ggc  
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Arg Leu Phe Cys Arg Thr Gln Trp Tyr Leu Arg Ile Asp Lys Arg Gly  
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Lys Val Lys Gly Thr Gln Glu Met Arg Asn Ser Tyr Asn Ile Met Glu  
85 90 95

20 atc atg act gtg gca gtt gga att gtg gca atc aaa ggg gtg gaa agt  
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Ile Met Thr Val Ala Val Gly Ile Val Ala Ile Lys Gly Val Glu Ser  
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25 gaa tac tat ctt gcc atg aac aaa caa ggg gaa ctc tat gca aag aaa  
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Glu Tyr Tyr Leu Ala Met Asn Lys Gln Gly Glu Leu Tyr Ala Lys Lys  
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30 gaa tgc aat gag gat tgc aac ttc aaa gaa ctg att ctg gaa aac cat  
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35 tac aac acc tct gca tca gct aaa tgg aca cac agc gga ggg gaa atg  
597  
Tyr Asn Thr Ser Ala Ser Ala Lys Trp Thr His Ser Gly Gly Glu Met  
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40 ttc gtg gcc tta aat caa aag ggg ctt cct gtc aaa ggg aag aaa acg  
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Phe Val Ala Leu Asn Gln Lys Gly Leu Pro Val Lys Gly Lys Lys Thr  
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45 aag aaa gaa caa aaa acg gcc cac ttt ctt cct atg gca ata act taa  
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tat gag cgc gat gaa ggt gat aaa tgg cga aac aaa aag ttt gaa ttg  
144  
Tyr Glu Arg Asp Glu Gly Asp Lys Trp Arg Asn Lys Lys Phe Glu Leu  
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192  
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240  
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288  
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Gly Ala Val Leu Asp Ile Arg Tyr Gly Val Ser Arg Ile Ala Tyr Ser  
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624

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20 gac tac atg gaa gga ggg gat ata agg gtg agg aga ctg ttc tgt cgc  
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864  
Thr Gln Trp Tyr Leu Arg Ile Asp Lys Arg Gly Lys Val Lys Gly Thr  
275 280 285

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Gln Glu Met Arg Asn Ser Tyr Asn Ile Met Glu Ile Met Thr Val Ala  
290 295 300

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40 atg aac aaa caa ggg gaa ctc tat gca aag aaa gaa tgc aat gag gat  
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1104  
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55 caa aag ggg ctt cct gtc aaa ggg aag aaa acg aag aaa gaa caa aaa  
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35 40 45

25 Gly Leu Glu Phe Pro Asn Leu Pro Tyr Tyr Ile Asp Gly Asp Val Lys
50 55 60

Leu Thr Gln Ser Met Ala Ile Ile Arg Tyr Ile Ala Asp Lys His Asn
30 65 70 75 80

Met Leu Gly Gly Cys Pro Lys Glu Arg Ala Glu Ile Ser Met Leu Glu
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35 Gly Ala Val Leu Asp Ile Arg Tyr Gly Val Ser Arg Ile Ala Tyr Ser
100 105 110

Lys Asp Phe Glu Thr Leu Lys Val Asp Phe Leu Ser Lys Leu Pro Glu
115 120 125

40 Met Leu Lys Met Phe Glu Asp Arg Leu Cys His Lys Thr Tyr Leu Asn
130 135 140

Gly Asp His Val Thr His Pro Asp Phe Met Leu Tyr Asp Ala Leu Asp
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Val Val Leu Tyr Met Asp Pro Met Cys Leu Asp Ala Phe Pro Lys Leu
165 170 175

50 Val Cys Phe Lys Lys Arg Ile Glu Ala Ile Pro Gln Ile Asp Lys Tyr
180 185 190

Leu Lys Ser Ser Lys Tyr Ile Ala Trp Pro Leu Gln Gly Trp Gln Ala
195 200 205

55 Thr Phe Gly Gly Gly Asp His Pro Pro Lys Ser Asp Leu Val Pro Arg
210 215 220

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